10/584960

SEQUENCE LISTING AP20 REC'D PCT/PTQ 30 JUN 2006

<110>	The University of North Carolina Threadgill, David W Lee, Daekee	
<120>	MODULATION OF EPIDERMAL GROWTH FACTOR HETERODIMER ACTIVITY	
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- Glu Ala Thr Cys Lys Asp Thr Cys Pro Pro Leu Met Leu Tyr Asn Pro Thr Thr Tyr Gln Met Asp Val Asn Pro Glu Gly Lys Tyr Ser Phe Gly 280 Ala Thr Cys Val Lys Lys Cys Pro Arg Asn Tyr Val Val Thr Asp His Gly Ser Cys Val Arg Ala Cys Gly Pro Asp Tyr Tyr Glu Val Glu Glu 310 315 Asp Gly Ile Arg Lys Cys Lys Cys Asp Gly Pro Cys Arg Lys Val 330 Cys Asn Gly Ile Gly Ile Gly Glu Phe Lys Asp Thr Leu Ser Ile Asn Ala Thr Asn Ile Lys His Phe Lys Tyr Cys Thr Ala Ile Ser Gly Asp 360 Leu His Ile Leu Pro Val Ala Phe Lys Gly Asp Ser Phe Thr Arg Thr 375 Pro Pro Leu Asp Pro Arg Glu Leu Glu Ile Leu Lys Thr Val Lys Glu 390 385 395 Ile Thr Gly Phe Leu Leu Ile Gln Ala Trp Pro Asp Asn Trp Thr Asp 405 410
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atg ga Met Gl 1															286
ccc cc Pro Pr															334

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				atc Ile								526
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-			_	cca Pro				 _	_	_		670
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				tgc Cys									1294
				agt Ser									1342
				ctg Leu									1390
				gcc Ala 390									1438
				atc Ile									1486
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				aat Asn									1582
				gly ggg									1630
				cat His 470									1678
				ttt Phe									1726
				gac Asp									1774

							tgc Cys 520									1822
gtc Val	aac Asn 530	tgc Cys	agc Ser	cag Gln	ttc Phe	ctt Leu 535	cgg Arg	ggc Gly	cag Gln	gag Glu	tgc Cys 540	gtg Val	gag Glu	gaa Glu	tgc Cys	1870
cga Arg 545	gta Val	ctg Leu	cag Gln	G] À aaa	ctc Leu 550	ccc Pro	agg Arg	gag Glu	tat Tyr	gtg Val 555	aat Asn	gcc Ala	agg Arg	cac His	tgt Cys 560	1918
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							gag Glu									2494

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cgg Arg					cag Gln		-	-	 _		_			•	2974
					ttt Phe										3022
					ctg Leu										3070
					gat Asp 950										3118
					cgg Arg										3166
					gac Asp										3214

	Leu			ccc tt Pro Le 10			Phe Ty		ege t Arg S		3262
				ggg Gly 1015							3307
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				aca Thr 1060							3442
				ctg Leu 1075							3487
				ctg Leu 1090							3532
	agc Ser 1100			gac Asp 1105					tac Tyr		3577
				ctg Leu 1120							3622
	ccc Pro 1130	Leu		ccc Pro 1135							3667
				cct Pro 1150							3712
				gcc Ala 1165							3757
				gtc Val 1180							3802
				gag Glu 1195							3847
		Gln		cct Pro 1210	Pro			Ala			3892

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ccc agc acc ttc aaa ggg aca cct acg gca gag aac cca gag tac Pro Ser Thr Phe Lys Gly Thr Pro Thr Ala Glu Asn Pro Glu Tyr 1235 1240 1245	3982
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Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His 35 40 45

Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr 50 60

Leu Pro Thr Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val 65 70 75 80

Gln Gly Tyr Val Leu Ile Ala His Asn Gln Val Arg Gln Val Pro Leu Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Asn Tyr Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Asn Asn Thr Thr Pro 120 Val Thr Gly Ala Ser Pro Gly Gly Leu Arg Glu Leu Gln Leu Arg Ser 135 Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Gln Arg Asn Pro Gln 155 Leu Cys Tyr Gln Asp Thr Ile Leu Trp Lys Asp Ile Phe His Lys Asn Asn Gln Leu Ala Leu Thr Leu Ile Asp Thr Asn Arg Ser Arg Ala Cys His Pro Cys Ser Pro Met Cys Lys Gly Ser Arg Cys Trp Gly Glu Ser Ser Glu Asp Cys Gln Ser Leu Thr Arg Thr Val Cys Ala Gly Gly Cys 215 220 Ala Arg Cys Lys Gly Pro Leu Pro Thr Asp Cys Cys His Glu Gln Cys 225 230 235 Ala Ala Gly Cys Thr Gly Pro Lys His Ser Asp Cys Leu Ala Cys Leu 245 250 His Phe Asn His Ser Gly Ile Cys Glu Leu His Cys Pro Ala Leu Val 260 265 270 Thr Tyr Asn Thr Asp Thr Phe Glu Ser Met Pro Asn Pro Glu Gly Arg 275 280 285 Tyr Thr Phe Gly Ala Ser Cys Val Thr Ala Cys Pro Tyr Asn Tyr Leu 290 295

315

Ser Thr Asp Val Gly Ser Cys Thr Leu Val Cys Pro Leu His Asn Gln

Glu Val Thr Ala Glu Asp Gly Thr Gln Arg Cys Glu Lys Cys Ser Lys Pro Cys Ala Arg Val Cys Tyr Gly Leu Gly Met Glu His Leu Arg Glu Val Arg Ala Val Thr Ser Ala Asn Ile Gln Glu Phe Ala Gly Cys Lys Lys Ile Phe Gly Ser Leu Ala Phe Leu Pro Glu Ser Phe Asp Gly Asp Pro Ala Ser Asn Thr Ala Pro Leu Gln Pro Glu Gln Leu Gln Val Phe Glu Thr Leu Glu Glu Ile Thr Gly Tyr Leu Tyr Ile Ser Ala Trp Pro Asp Ser Leu Pro Asp Leu Ser Val Phe Gln Asn Leu Gln Val Ile Arg Gly Arg Ile Leu His Asn Gly Ala Tyr Ser Leu Thr Leu Gln Gly Leu Gly Ile Ser Trp Leu Gly Leu Arg Ser Leu Arg Glu Leu Gly Ser Gly Leu Ala Leu Ile His His Asn Thr His Leu Cys Phe Val His Thr Val Pro Trp Asp Gln Leu Phe Arg Asn Pro His Gln Ala Leu Leu His Thr Ala Asn Arg Pro Glu Asp Glu Cys Val Gly Glu Gly Leu Ala Cys His Gln Leu Cys Ala Arg Gly His Cys Trp Gly Pro Gly Pro Thr Gln Cys Val Asn Cys Ser Gln Phe Leu Arg Gly Gln Glu Cys Val Glu Glu Cys

Arg Val Leu Gln Gly Leu Pro Arg Glu Tyr Val Asn Ala Arg His Cys

Leu Pro Cys His Pro Glu Cys Gln Pro Gln Asn Gly Ser Val Thr Cys Phe Gly Pro Glu Ala Asp Gln Cys Val Ala Cys Ala His Tyr Lys Asp Pro Pro Phe Cys Val Ala Arg Cys Pro Ser Gly Val Lys Pro Asp Leu Ser Tyr Met Pro Ile Trp Lys Phe Pro Asp Glu Glu Gly Ala Cys Gln Pro Cys Pro Ile Asn Cys Thr His Ser Cys Val Asp Leu Asp Asp Lys Gly Cys Pro Ala Glu Gln Arg Ala Ser Pro Leu Thr Ser Ile Ile Ser Ala Val Val Gly Ile Leu Leu Val Val Leu Gly Val Val Phe Gly Ile Leu Ile Lys Arg Arg Gln Gln Lys Ile Arg Lys Tyr Thr Met Arg Arg Leu Leu Gln Glu Thr Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Ala Met Pro Asn Gln Ala Gln Met Arg Ile Leu Lys Glu Thr Glu Leu Arg Lys Val Lys Val Leu Gly Ser Gly Ala Phe Gly Thr Val Tyr Lys Gly Ile Trp Ile Pro Asp Gly Glu Asn Val Lys Ile Pro Val Ala Ile Lys Val Leu Arg Glu Asn Thr Ser Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Gly Val Gly Ser Pro Tyr Val Ser Arg

Leu Leu Gly Ile Cys Leu Thr Ser Thr Val Gln Leu Val Thr Gln Leu

- Met Pro Tyr Gly Cys Leu Leu Asp His Val Arg Glu Asn Arg Gly Arg 805 810 815
- Leu Gly Ser Gln Asp Leu Leu Asn Trp Cys Met Gln Ile Ala Lys Gly 820 825 830
- Met Ser Tyr Leu Glu Asp Val Arg Leu Val His Arg Asp Leu Ala Ala 835 840 845
- Arg Asn Val Leu Val Lys Ser Pro Asn His Val Lys Ile Thr Asp Phe 850 855 860
- Gly Leu Ala Arg Leu Leu Asp Ile Asp Glu Thr Glu Tyr His Ala Asp 865 870 875 880
- Gly Gly Lys Val Pro Ile Lys Trp Met Ala Leu Glu Ser Ile Leu Arg 885 890 895
- Arg Arg Phe Thr His Gln Ser Asp Val Trp Ser Tyr Gly Val Thr Val 900 905 910
- Trp Glu Leu Met Thr Phe Gly Ala Lys Pro Tyr Asp Gly Ile Pro Ala 915 920 925
- Arg Glu Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg Leu Pro Gln Pro 930 935 940
- Pro Ile Cys Thr Ile Asp Val Tyr Met Ile Met Val Lys Cys Trp Met 945 950 955 960
- Ile Asp Ser Glu Cys Arg Pro Arg Phe Arg Glu Leu Val Ser Glu Phe 965 970 975
- Ser Arg Met Ala Arg Asp Pro Gln Arg Phe Val Val Ile Gln Asn Glu 980 985 990
- Asp Leu Gly Pro Ala Ser Pro Leu Asp Ser Thr Phe Tyr Arg Ser Leu 995 1000 1005
- Leu Glu Asp Asp Asp Met Gly Asp Leu Val Asp Ala Glu Glu Tyr 1010 1015 1020
- Leu Val Pro Gln Gln Gly Phe Phe Cys Pro Asp Pro Ala Pro Gly 1025 1030 1035

Ala Gly Gly Met Val His His Arg His Arg Ser Ser Ser Thr Arg Ser Gly Gly Gly Asp Leu Thr Leu Gly Leu Glu Pro Ser Glu Glu Glu Ala Pro Arg Ser Pro Leu Ala Pro Ser Glu Gly Ala Gly Ser Asp Val Phe Asp Gly Asp Leu Gly Met Gly Ala Ala Lys Gly Leu Gln Ser Leu Pro Thr His Asp Pro Ser Pro Leu Gln Arg Tyr Ser Glu Asp Pro Thr Val Pro Leu Pro Ser Glu Thr Asp Gly Tyr Val Ala Pro Leu Thr Cys Ser Pro Gln Pro Glu Tyr Val Asn Gln Pro Asp Val Arg Pro Gln Pro Pro Ser Pro Arg Glu Gly Pro Leu Pro Ala Ala Arg Pro Ala Gly Ala Thr Leu Glu Arg Pro Lys Thr Leu Ser Pro Gly Lys Asn Gly Val Val Lys Asp Val Phe Ala Phe Gly Gly Ala Val Glu Asn Pro Glu Tyr Leu Thr Pro Gln Gly Gly Ala Ala Pro Gln Pro His Pro Pro Pro Ala Phe Ser Pro Ala Phe Asp Asn Leu Tyr Tyr Trp Asp Gln Asp Pro Pro Glu Arg Gly Ala Pro Pro Ser Thr Phe Lys Gly Thr Pro Thr Ala Glu Asn Pro Glu Tyr Leu Gly Leu Asp Val Pro Val

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atq gag ctg gcg tqg tqc cqt tqg qqg ttc ctc ctc qcc ctc ctg
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Met Glu Leu Ala Ala Trp Cys Arg Trp Gly Phe Leu Leu Ala Leu Leu
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tcc ccc gga gcc gcg ggt acc caa gtg tgt acc ggt acc gac atg aag
Ser Pro Gly Ala Ala Gly Thr Gln Val Cys Thr Gly Thr Asp Met Lys
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ttg cga ctc cct gcc agt cct gag acc cac ctg gac atg ctt cgc cac
Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His
        35
ctc tac cag ggc tgt cag gtg cag ggc aat ttg gag ctt acc tac
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Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr
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ctg ccc gcc aat gcc agc ctc tca ttc ctg cag gac atc cag gaa gtc
                                                                     419
Leu Pro Ala Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val
65
                    70
                                                            80
                                                                      467
cag gga tac atg ctc atc gct cac aac cga gtg aaa cac gtc cca ctg
Gln Gly Tyr Met Leu Ile Ala His Asn Arg Val Lys His Val Pro Leu
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cag agg ttg cgc atc gtg aga ggg act cag ctc ttt gag gac aag tat
Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Lys Tyr
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                                105
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gcc ctg gct gtg cta gac aac cga gac cct ttg gac aac gtc acc acc
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Ala Leu Ala Val Leu Asp Asn Arg Asp Pro Leu Asp Asn Val Thr Thr
        115
                            120
gcc gcc cca ggc aga acc cca gaa ggg ctg cgg gag ctg cag ctt cga
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Ala Ala Pro Gly Arg Thr Pro Glu Gly Leu Arg Glu Leu Gln Leu Arg
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                        135
                                                                     659
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Ser Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Arg Gly Asn Pro
145
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							tgc Cys 300			1091
	_	-	 	_	_	_	tgt Cys	_		1139
							tgt Cys			1187
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							gag Glu 380			1331
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					cat His							1	.523
_					ctg Leu		_	_		 _	 -	1	.571
	_	-			cac His 470	_			_	_		1	619
					ctc Leu							1	.667
					gaa Glu							1	715
		_	-	-	cgt Arg		_	 				1	.763
					cag Gln							1	.811
					ggg Gly 550							1	.859
					ccc Pro							1	.907
					gct Ala							1	955
					gtg Val							2	2003
					atc Ile							2	2051
					aac Asn 630					_		2	2099

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				ctg Leu							2195
				agg Arg							2243
				acc Thr 695							2291
				gct Ala							2339
				ctt Leu							2387
				gat Asp							2435
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Ser Pro Gly Ala Ala Gly Thr Gln Val Cys Thr Gly Thr Asp Met Lys 20 25 30

Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His $35 \hspace{1cm} 40 \hspace{1cm} 45$

Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr 50 55 60

Leu Pro Ala Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val 65 70 75 80

Gln Gly Tyr Met Leu Ile Ala His Asn Arg Val Lys His Val Pro Leu $85 \hspace{1cm} 90 \hspace{1cm} 95$

Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Lys Tyr 100 105 110

Ala Leu Ala Val Leu Asp Asn Arg Asp Pro Leu Asp Asn Val Thr Thr 115 120 125

Ala Ala Pro Gly Arg Thr Pro Glu Gly Leu Arg Glu Leu Gln Leu Arg 130 135 140

Ser Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Arg Gly Asn Pro Gln Leu Cys Tyr Gln Asp Met Val Leu Trp Lys Asp Val Leu Arg Lys Asn Asn Gln Leu Ala Pro Val Asp Met Asp Thr Asn Arg Ser Arg Ala Cys Pro Pro Cys Ala Pro Thr Cys Lys Asp Asn His Cys Trp Gly Glu Ser Pro Glu Asp Cys Gln Ile Leu Thr Gly Thr Ile Cys Thr Ser Gly Cys Ala Arg Cys Lys Gly Arg Leu Pro Thr Asp Cys Cys His Glu Gln Cys Ala Ala Gly Cys Thr Gly Pro Lys His Ser Asp Cys Leu Ala Cys Leu His Phe Asn His Ser Gly Ile Cys Glu Leu His Cys Pro Ala Leu Ile Thr Tyr Asn Thr Asp Thr Phe Glu Ser Met Leu Asn Pro Glu Gly Arg Tyr Thr Phe Gly Ala Ser Cys Val Thr Thr Cys Pro Tyr Asn Tyr Leu Ser Thr Glu Val Gly Ser Cys Thr Leu Val Cys Pro Pro Asn Asn Gln Glu Val Thr Ala Glu Asp Gly Thr Gln Arg Cys Glu Lys Cys Ser Lys Pro Cys Ala Gly Val Cys Tyr Gly Leu Gly Met Glu His Leu Arg Gly Ala Arg Ala Ile Thr Ser Asp Asn Ile Gln Glu Phe Ala Gly Cys Lys Lys Ile Phe Gly Ser Leu Ala Phe Leu Pro Glu Ser Phe Asp Gly

Asn 385	Pro	Ser	Ser	Gly	Val 390	Ala	Pro	Leu	Lys	Pro 395	Glu	His	Leu	Gln	Val 400
Phe	Glu	Thr	Leu	Glu 405	Glu	Ile	Thr	Gly	Tyr 410	Leu	Tyr	Ile	Ser	Ala 415	Trp
Pro	Glu	Ser	Phe 420	Gln	Asp	Leu	Ser	Val 425	Phe	Gln	Asn	Leu	Arg 430	Val	Ile
Arg	Gly	Arg 435	Ile	Leu	His	Asp	Gly 440	Ala	Tyr	Ser	Leu	Thr 445	Leu	Gln	Gly
Leu	Gly 450	Ile	His	Ser	Leu	Gly 455	Leu	Arg	Ser	Leu	Arg 460	Glu	Leu	Gly	Ser
Gly 465	Leu	Ala	Leu	Ile	His 470	Arg	Asn	Thr	His	Leu 475	Cys	Phe	Val	Asn	Thr 480
Val	Pro	Trp	Asp	Gln 485	Leu	Phe	Arg	Asn	Pro 490	His	Gln	Ala	Leu	Leu 495	His
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Asn	Ser	Leu 515	Cys	Ala	Arg	Gly	His 520	Cys	Trp	Gly	Pro	Gly 525	Pro	Thr	Gln
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Cys 545	Arg	Val	Trp	Lys	Gly 550	Leu	Pro	Arg	Glu	Tyr 555	Val	Arg	Gly	Lys	His 560
Суѕ	Leu	Pro	Суѕ	His 565	Pro	Glu	Cys	Gln	Pro 570	Gln	Asn	Ser	Ser	Glu 575	Thr
Cys	Tyr	Gly	Ser 580	Glu	Ala	Asp	Gln	Cys 585	Glu	Ala	Cys	Ala	His 590	Tyr	Lys
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Leu	Ser 610	Tyr	Met	Pro	Ile	Trp 615	Lys	Tyr	Pro	Asp	Glu 620	Glu	Gly	Ile	Cys

Gln Pro Cys Pro Ile Asn Cys Thr His Ser Cys Val Asp Leu Asp Glu Arg Gly Cys Pro Ala Glu Gln Arg Ala Ser Pro Val Thr Phe Ile Ile 645 . Ala Thr Val Val Gly Val Leu Leu Phe Leu Ile Ile Val Val Val Ile Gly Ile Leu Ile Lys Arg Arg Gln Lys Ile Arg Lys Tyr Thr Met Arg Arg Leu Leu Gln Glu Thr Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Ala Val Pro Asn Gln Ala Gln Met Arg Ile Leu Lys Glu Thr Glu Leu Arg Lys Leu Lys Val Leu Gly Ser Gly Ala Phe Gly Thr Val Tyr Lys Gly Ile Trp Ile Pro Asp Gly Glu Asn Val Lys Ile Pro Val Ala Ile Lys Val Leu Arg Glu Asn Thr Ser Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Gly Val Gly Ser Pro Tyr Val Ser Arg Leu Leu Gly Ile Cys Leu Thr Ser Thr Val Gln Leu Val Thr Gln Leu Met Pro Tyr Gly Cys Leu Leu Asp His Val Arg Glu His Arg Gly Arg Leu Gly Ser Gln Asp Leu Leu Asn Trp Cys Val Gln Ile Ala Lys Gly Met Ser Tyr Leu Glu Glu Val Arg Leu Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Lys Ser Pro Asn His Val Lys Ile Thr Asp

- Phe Gly Leu Ala Arg Leu Leu Asp Ile Asp Glu Thr Glu Tyr His Ala 865 870 875 880
- Asp Gly Gly Lys Val Pro Ile Lys Trp Met Ala Leu Glu Ser Ile Leu 885 890 895
- Arg Arg Arg Phe Thr His Gln Ser Asp Val Trp Ser Tyr Gly Val Thr 900 905 910
- Val Trp Glu Leu Met Thr Phe Gly Ala Lys Pro Tyr Asp Gly Ile Pro 915 920 925
- Ala Arg Glu Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg Leu Pro Gln 930 935 940
- Pro Pro Ile Cys Thr Ile Asp Val Tyr Met Ile Met Val Lys Cys Trp 945 950 955 960
- Met Ile Asp Ser Glu Cys Arg Pro Arg Phe Arg Glu Leu Val Ser Glu 965 970 975
- Phe Ser Arg Met Ala Arg Asp Pro Gln Arg Phe Val Val Ile Gln Asn 980 985 990
- Glu Asp Leu Gly Pro Ser Ser Pro Met Asp Ser Thr Phe Tyr Arg Ser 995 1000 1005
- Leu Leu Glu Asp Asp Asp Met Gly Glu Leu Val Asp Ala Glu Glu 1010 1015 1020
- Tyr Leu Val Pro Gl
n Gl
n Gly Phe Phe Ser Pro Asp Pro Ala Leu 1025 1030 1035
- Gly Thr Gly Ser Thr Ala His Arg Arg His Arg Ser Ser Ser Ala 1040 1045 1050
- Arg Ser Gly Gly Glu Leu Thr Leu Gly Leu Glu Pro Ser Glu 1055 1060 1065
- Glu Glu Pro Pro Arg Ser Pro Leu Ala Pro Ser Glu Gly Ala Gly 1070 1080
- Ser Asp Val Phe Asp Gly Asp Leu Ala Val Gly Val Thr Lys Gly 1085 1090 1095

Leu Gln Ser Leu Ser Pro His Asp Leu Ser Pro Leu Gln Arg Tyr 1100 1105 Ser Glu Asp Pro Thr Leu Pro Leu Pro Pro Glu Thr Asp Gly Tyr 1120 1115 Val Ala Pro Leu Ala Cys Ser Pro Gln Pro Glu Tyr Val Asn Gln 1135 Pro Glu Val Arg Pro Gln Ser Pro Leu Thr Pro Glu Gly Pro Pro 1150 Pro Pro Ile Arg Pro Ala Gly Ala Thr Leu Glu Arg Pro Lys Thr 1165 Leu Ser Pro Gly Lys Asn Gly Val Val Lys Asp Val Phe Ala Phe 1175 1180 Gly Gly Ala Val Glu Asn Pro Glu Tyr Leu Ala Pro Arg Ala Gly 1190 1195 Thr Ala Ser Gln Pro His Pro Ser Pro Ala Phe Ser Pro Ala Phe 1205 1210 1215 Asp Asn Leu Tyr Tyr Trp Asp Gln Asn Ser Ser Glu Gln Gly Pro 1220 1225 1230 Pro Pro Ser Thr Phe Glu Gly Thr Pro Thr Ala Glu Asn Pro Glu 1235 1240 1245 Tyr Leu Gly Leu Asp Val Pro Val 1250 <210> 14 <211> 5511 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (194)..(4222) <400> 14 acacacaca accectecee tgecatecet ecceggaete eggeteegge teegattgea atttgcaacc teegetgeeg tegeegeage agecaceaat tegeeagegg tteaggtgge

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tgt cct ggg act ctg Cys Pro Gly Thr Leu 30			
caa tac cag aca ctg Gln Tyr Gln Thr Leu 45			
ggg aac ctt gag att Gly Asn Leu Glu Ile 65		= =	
ctg cag tgg att cga Leu Gln Trp Ile Arg 80			
gaa ttc tct act cta Glu Phe Ser Thr Leu 95	=		= = =
cag gtc tac gat ggg Gln Val Tyr Asp Gly 110			
acc aac tcc agc cac Thr Asn Ser Ser His 125			
gag att ctg tca ggg Glu Ile Leu Ser Gly 145			
cac atg gac aca att His Met Asp Thr Ile 160			g Asp Ala
gag ata gtg gtg aag Glu Ile Val Val Lys 175			
gtt tgc aag ggg cga Val Cys Lys Gly Arg 190			
ttg acc aag acc atc Leu Thr Lys Thr Ile 205			

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					tgc Cys							949
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					gga Gly							1477
					aac Asn 435							1525
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					cac His										1621
	_	_	, ,	_	cta Leu	-		_			_	_	_	_	1669
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_			_	_	gtg Val 530			_		_					1813
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					aaa Lys										2245
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	_		_	_	gct Ala	_	-				_		2725
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					aca Thr 900								2917
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acc ttt aaa gaa cta gcc aat gag ttc acc agg atg gcc cga gac cca Thr Phe Lys Glu Leu Ala Asn Glu Phe Thr Arg Met Ala Arg Asp Pro 960 965 970	3109
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ggg cca gag ccc cat ggt ctg aca aac aag aag cta gag gaa gta gag Gly Pro Glu Pro His Gly Leu Thr Asn Lys Lys Leu Glu Glu Val Glu 990 995 1000	3205
ctg gag cca gaa cta gac cta gac cta gac ttg gaa gca gag gag Leu Glu Pro Glu Leu Asp Leu Asp Leu Asp Leu Glu Ala Glu Glu 1005 1010 1015	3250
gac aac ctg gca acc acc aca ctg ggc tcc gcc ctc agc cta cca Asp Asn Leu Ala Thr Thr Thr Leu Gly Ser Ala Leu Ser Leu Pro 1020 1025 1030	3295
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cca tca tct gga tac atg ccc atg aac cag ggt aat ctt ggg gag Pro Ser Ser Gly Tyr Met Pro Met Asn Gln Gly Asn Leu Gly Glu 1050 1055 1060	3385
tct tgc cag gag tct gca gtt tct ggg agc agt gaa cgg tgc ccc Ser Cys Gln Glu Ser Ala Val Ser Gly Ser Ser Glu Arg Cys Pro 1065 1070 1075	3430
cgt cca gtc tct cta cac cca atg cca cgg gga tgc ctg gca tca Arg Pro Val Ser Leu His Pro Met Pro Arg Gly Cys Leu Ala Ser 1080 1085 1090	3475
gag toa toa gag ggg cat gta aca ggc tot gag got gag oto cag Glu Ser Ser Glu Gly His Val Thr Gly Ser Glu Ala Glu Leu Gln 1095 1100 1105	3520
gag aaa gtg tca atg tgt agg agc cgg agc agg agc cgg agc cca Glu Lys Val Ser Met Cys Arg Ser Arg Ser Arg Ser Arg Ser Pro 1110 1115 1120	3565
cgg cca cgc gga gat agc gcc tac cat tcc cag cgc cac agt ctg Arg Pro Arg Gly Asp Ser Ala Tyr His Ser Gln Arg His Ser Leu 1125 1130 1135	3610
ctg act cct gtt acc cca ctc tcc cca ccc ggg tta gag gaa gag Leu Thr Pro Val Thr Pro Leu Ser Pro Pro Gly Leu Glu Glu 1140 1145 1150	3655
gat gtc aac ggt tat gtc atg cca gat aca cac ctc aaa ggt act Asp Val Asn Gly Tyr Val Met Pro Asp Thr His Leu Lys Gly Thr 1155 1160 1165	3700

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gtc Val 1185	Leu				gaa Glu 1190										3790
atg Met 1200	Asn			_	agg Arg 1205		_					cct Pro			3835
agt Ser 1215	Ser				ctg Leu 1220										3880
gac Asp 1230	Leu				ctg Leu 1235										3925
cct Pro 1245	_			_	ccc Pro 1250		_					_	_	-	3970
tat Tyr 1260	Ğlu				cgg Arg 1265		-	-							4015
gat Asp 1275	Tyr	-	-	_	ggg Gly 1280	_	_		-						4060
gaa Glu 1290		_	_	_	ttt Phe 1295	_					_	-			4105
gtc Val 1305	His		_	_	cta Leu 1310				_	-		gag Glu	-		4150
gac Asp 1320					aac Asn 1325										4195
ccc Pro 1335	Lys	-		_	cag Gln 1340	_	_	taa	ctc	ctgct	cc ct	tgtg	gcact	t	4242
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tctc	tccc	ag g	tccc	agcc	c ctt	ttcc	cca (gtcc	cagad	ca at	tcca	ttca	atct	tttggag	4362
gctt	ttaa	ac a	tttt	gacad	c aaaa	attci	tta 1	tggta	atgta	ag cca	agcto	gtgc	actt	ttcttct	4422
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gctt	cttca	ac a	ggca	ctcct	gga	gata	tga a	aggat	ttact	c to	cata	tccc	ttc	ctctcag	4542
gctc	ttga	ct a	cttg	gaact	aggo	ctcti	tat o	gtgt	gcctt	t gt	ttcc	catc	agad	ctgtcaa	4602

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<212> PRT

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Ala Arg Gly Ser Glu Val Gly Asn Ser Gln Ala Val Cys Pro Gly Thr 20 25 30

Leu Asn Gly Leu Ser Val Thr Gly Asp Ala Glu Asn Gln Tyr Gln Thr 35 40 45

Leu Tyr Lys Leu Tyr Glu Arg Cys Glu Val Val Met Gly Asn Leu Glu 50 60

Ile Val Leu Thr Gly His Asn Ala Asp Leu Ser Phe Leu Gln Trp Ile 70 75 80

- Arg Glu Val Thr Gly Tyr Val Leu Val Ala Met Asn Glu Phe Ser Thr 85 90 95
- Leu Pro Leu Pro Asn Leu Arg Val Val Arg Gly Thr Gln Val Tyr Asp $100 \hspace{1.5cm} 105 \hspace{1.5cm} 110 \hspace{1.5cm}$
- Gly Lys Phe Ala Ile Phe Val Met Leu Asn Tyr Asn Thr Asn Ser Ser 115 120 125
- His Ala Leu Arg Gln Leu Arg Leu Thr Gln Leu Thr Glu Ile Leu Ser 130 135 140 .
- Gly Gly Val Tyr Ile Glu Lys Asn Asp Lys Leu Cys His Met Asp Thr 145 150 155 160
- Ile Asp Trp Arg Asp Ile Val Arg Asp Arg Asp Ala Glu Ile Val Val 165 170 175
- Lys Asp Asn Gly Arg Ser Cys Pro Pro Cys His Glu Val Cys Lys Gly 180 185 190
- Arg Cys Trp Gly Pro Gly Ser Glu Asp Cys Gln Thr Leu Thr Lys Thr 195 200 205
- Ile Cys Ala Pro Gln Cys Asn Gly His Cys Phe Gly Pro Asn Pro Asn 210 215 220
- Gln Cys Cys His Asp Glu Cys Ala Gly Gly Cys Ser Gly Pro Gln Asp 225 230 235 240
- Thr Asp Cys Phe Ala Cys Arg His Phe Asn Asp Ser Gly Ala Cys Val 245 250 255
- Pro Arg Cys Pro Gln Pro Leu Val Tyr Asn Lys Leu Thr Phe Gln Leu 260 265 270
- Glu Pro Asn Pro His Thr Lys Tyr Gln Tyr Gly Gly Val Cys Val Ala 275 280 285
- Ser Cys Pro His Asn Phe Val Val Asp Gln Thr Ser Cys Val Arg Ala 290 295 300
- Cys Pro Pro Asp Lys Met Glu Val Asp Lys Asn Gly Leu Lys Met Cys 305 310 315 320

- Glu Pro Cys Gly Gly Leu Cys Pro Lys Ala Cys Glu Gly Thr Gly Ser 325 330 335
- Gly Ser Arg Phe Gln Thr Val Asp Ser Ser Asn Ile Asp Gly Phe Val 340 345 . 350
- Asn Cys Thr Lys Ile Leu Gly Asn Leu Asp Phe Leu Ile Thr Gly Leu 355 360 365
- Asn Gly Asp Pro Trp His Lys Ile Pro Ala Leu Asp Pro Glu Lys Leu 370 380
- Asn Val Phe Arg Thr Val Arg Glu Ile Thr Gly Tyr Leu Asn Ile Gln 385 390 395 400
- Ser Trp Pro Pro His Met His Asn Phe Ser Val Phe Ser Asn Leu Thr 405 410 415
- Thr Ile Gly Gly Arg Ser Leu Tyr Asn Arg Gly Phe Ser Leu Leu Ile 420 425 430
- Met Lys Asn Leu Asn Val Thr Ser Leu Gly Phe Arg Ser Leu Lys Glu 435 440 445
- Ile Ser Ala Gly Arg Ile Tyr Ile Ser Ala Asn Arg Gln Leu Cys Tyr 450 460
- His His Ser Leu Asn Trp Thr Lys Val Leu Arg Gly Pro Thr Glu Glu 465 470 475 480
- Arg Leu Asp Ile Lys His Asn Arg Pro Arg Arg Asp Cys Val Ala Glu 485 490 495
- Gly Lys Val Cys Asp Pro Leu Cys Ser Ser Gly Gly Cys Trp Gly Pro 500 505 510
- Gly Pro Gly Gln Cys Leu Ser Cys Arg Asn Tyr Ser Arg Gly Gly Val 515 520 525
- Cys Val Thr His Cys Asn Phe Leu Asn Gly Glu Pro Arg Glu Phe Ala 530 540
- His Glu Ala Glu Cys Phe Ser Cys His Pro Glu Cys Gln Pro Met Glu 545 550 555 560

Gly Thr Ala Thr Cys Asn Gly Ser Gly Ser Asp Thr Cys Ala Gln Cys Ala His Phe Arg Asp Gly Pro His Cys Val Ser Ser Cys Pro His Gly Val Leu Gly Ala Lys Gly Pro Ile Tyr Lys Tyr Pro Asp Val Gln Asn Glu Cys Arg Pro Cys His Glu Asn Cys Thr Gln Gly Cys Lys Gly Pro Glu Leu Gln Asp Cys Leu Gly Gln Thr Leu Val Leu Ile Gly Lys Thr His Leu Thr Met Ala Leu Thr Val Ile Ala Gly Leu Val Val Ile Phe Met Met Leu Gly Gly Thr Phe Leu Tyr Trp Arg Gly Arg Arg Ile Gln Asn Lys Arg Ala Met Arg Arg Tyr Leu Glu Arg Gly Glu Ser Ile Glu Pro Leu Asp Pro Ser Glu Lys Ala Asn Lys Val Leu Ala Arg Ile Phe Lys Glu Thr Glu Leu Arg Lys Leu Lys Val Leu Gly Ser Gly Val Phe Gly Thr Val His Lys Gly Val Trp Ile Pro Glu Gly Glu Ser Ile Lys Ile Pro Val Cys Ile Lys Val Ile Glu Asp Lys Ser Gly Arg Gln Ser Phe Gln Ala Val Thr Asp His Met Leu Ala Ile Gly Ser Leu Asp His Ala His Ile Val Arg Leu Leu Gly Leu Cys Pro Gly Ser Ser Leu Gln

Leu Val Thr Gln Tyr Leu Pro Leu Gly Ser Leu Leu Asp His Val Arg

- Gln His Arg Gly Ala Leu Gly Pro Gln Leu Leu Leu Asn Trp Gly Val 805 810 815
- Gln Ile Ala Lys Gly Met Tyr Tyr Leu Glu Glu His Gly Met Val His 820 825 830
- Arg Asn Leu Ala Ala Arg Asn Val Leu Leu Lys Ser Pro Ser Gln Val 835 840 845
- Gln Val Ala Asp Phe Gly Val Ala Asp Leu Pro Pro Asp Asp Lys 850 855 860
- Gln Leu Leu Tyr Ser Glu Ala Lys Thr Pro Ile Lys Trp Met Ala Leu 865 870 875 880
- Glu Ser Ile His Phe Gly Lys Tyr Thr His Gln Ser Asp Val Trp Ser 885 890 895
- Tyr Gly Val Thr Val Trp Glu Leu Met Thr Phe Gly Ala Glu Pro Tyr 900 905 910
- Ala Gly Leu Arg Leu Ala Glu Val Pro Asp Leu Leu Glu Lys Gly Glu 915 920 925
- Arg Leu Ala Gln Pro Gln Ile Cys Thr Ile Asp Val Tyr Met Val Met 930 935 940
- Val Lys Cys Trp Met Ile Asp Glu Asn Ile Arg Pro Thr Phe Lys Glu 945 950 955 960
- Leu Ala Asn Glu Phe Thr Arg Met Ala Arg Asp Pro Pro Arg Tyr Leu 965 970 . 975
- Val Ile Lys Arg Glu Ser Gly Pro Gly Ile Ala Pro Gly Pro Glu Pro 980 985 990
- His Gly Leu Thr Asn Lys Lys Leu Glu Glu Val Glu Leu Glu Pro Glu 995 1000 1005
- Leu Asp Leu Asp Leu Glu Ala Glu Glu Asp Asn Leu Ala 1010 1015 1020
- Thr Thr Thr Leu Gly Ser Ala Leu Ser Leu Pro Val Gly Thr Leu 1025 . 1030 1035

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Tyr	Met 1055	Pro	Met	Asn	Gln	Gly 1060	Asn	Leu	Gly	Glu	Ser 1065	Cys	Gln	Glu
Ser	Ala 1070	Val	Ser	Gly	Ser	Ser 1075	Glu	Arg	Суѕ	Pro	Arg 1080	Pro	Val	Ser
Leu	His 1085	Pro	Met	Pro	Arg	Gly 1090	Cys	Leu	Ala	Ser	Glu 1095	Ser	Ser	Glu
Gly	His 1100	Val	Thr	Gly	Ser	Glu 1105		Glu	Leu	Gln	Glu 1110	Lys	Val	Ser
Met	Cys 1115	Arg	Ser	Arg	Ser	Arg 1120	Ser	Arg	Ser	Pro	Arg 1125	Pro	Arg	Gly
Asp	Ser 1130		Tyr	His	Ser	Gln 1135	Arg	His	Ser	Leu	Leu 1140	Thr	Pro	Val
Thr	Pro 1145	Leu	Ser	Pro	Pro	Gly 1150	Leu	Glu	Glu	Glu	Asp 1155		Asn	Gly
Tyr	Val 1160	Met	Pro	Asp	Thr	His 1165	Leu	Lys	Gly	Thr	Pro 1170	Ser	Ser	Arg
Glu	Gly 1175		Leu	Ser	Ser	Val 1180	_	Leu	Ser	Ser	Val 1185	Leu	Gly	Thr
			-		_	Glu 1195		_		_			Arg	Arg
Arg	Arg 1205	His	Ser	Pro	Pro	His 1210	Pro	Pro	Arg	Pro	Ser 1215	Ser	Leu	Glu
Glu	Leu 1220	Gly	Tyr	Glu	Tyr	Met 1225	Asp	Val	Gly	Ser	Asp 1230	Leu	Ser	Ala
Ser	Leu 1235	Gly	Ser	Thr	Gln	Ser 1240	Cys	Pro	Leu	His	Pro 1245	Val	Pro	Ile
Met	Pro 1250	Thr	Ala	Gly	Thr	Thr 1255	Pro	Asp	Glu	Asp	Tyr 1260	Glu	Tyr	Met

Met Gly Ala Cys Pro Ala Ser Glu Gln Gly Tyr Glu Glu Met Arg 1280 1285 1290	
Ala Phe Gln Gly Pro Gly His Gln Ala Pro His Val His Tyr Ala 1295 1300 1305	
Arg Leu Lys Thr Leu Arg Ser Leu Glu Ala Thr Asp Ser Ala Phe 1310 1315 1320	
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atg agt gcg att ggg act ctg cag gtg ctg ggt ttc ctt ctc agc ctg Met Ser Ala Ile Gly Thr Leu Gln Val Leu Gly Phe Leu Leu Ser Leu 1 5 10 15	48 96
atg agt gcg att ggg act ctg cag gtg ctg ggt ttc ctt ctc agc ctg Met Ser Ala Ile Gly Thr Leu Gln Val Leu Gly Phe Leu Leu Ser Leu 1 5 10 15 gcc cgg ggt tcc gag atg ggc aac tct cag gca gta tgt cct ggg act Ala Arg Gly Ser Glu Met Gly Asn Ser Gln Ala Val Cys Pro Gly Thr 20 25 30	
atg agt gcg att ggg act ctg cag gtg ctg ggt ttc ctt ctc agc ctg Met Ser Ala Ile Gly Thr Leu Gln Val Leu Gly Phe Leu Leu Ser Leu 1 5 10 15 gcc cgg ggt tcc gag atg ggc aac tct cag gca gta tgt cct ggg act Ala Arg Gly Ser Glu Met Gly Asn Ser Gln Ala Val Cys Pro Gly Thr 20 25 30 cta aac ggg ctg agt gtg acc ggc gat gct gac aac cag tac cag aca Leu Asn Gly Leu Ser Val Thr Gly Asp Ala Asp Asn Gln Tyr Gln Thr 35 40 45	96
atg agt gcg att ggg act ctg cag gtg ctg ggt ttc ctt ctc agc ctg Met Ser Ala Ile Gly Thr Leu Gln Val Leu Gly Phe Leu Leu Ser Leu 1	96 44

					ctc Leu											336
					ttt Phe											384
					ctc Leu											432
					gag Glu 150											480
att Ile	gac Asp	tgg Trp	agg Arg	gac Asp 165	atc Ile	gtg Val	agg Arg	gtt Val	cca Pro 170	gac Asp	gct Ala	gag Glu	ata Ile	gtg Val 175	gtg Val	528
					aac Asn											576
					gga Gly											624
					tgt Cys											672
					gaa Glu 230											720
					tgc Cys											768
ccc Pro	agg Arg	tgt Cys	cca Pro 260	gcg Ala	ccc Pro	ctt Leu	gtg Val	tac Tyr 265	aac Asn	aag Lys	cta Leu	acg Thr	ttc Phe 270	cag Gln	ctt Leu	816
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					ttt Phe											912
				Lys	atg Met 310											960
					ctg Leu											1008

					acc Thr											1056
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					cac His											1152
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	_	-		_	gtc Val			_	_		_			_		1392
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			_	_	cca Pro	_	_					_				1536
					ttg Leu											1584
					aac Asn											1632
					ttc Phe 550											1680
		_	_	_	aat Asn		_			_	-	_	_	_	_	1728

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			ggt Gly									182	:4
			cac His									187	2
			tta Leu 630									192	:0
			gtg Val									196	8
_	 		ctc Leu			_	 _			_		201	. 6
			tac Tyr									206	54
			gca Ala									211	.2
			ctt Leu 710									216	60
			tgg Trp									220	18
			atc Ile									225	6
_		-	atg Met	_	-	_	 _	_	_		_	230	14
			gga Gly									235	2
			ctg Leu 790									240	0
			cca Pro									244	8

	aag Lys															2496
	gcg Ala															2544
	gat Asp 850															2592
	cac His															2640
	cac His															2688
_	acc Thr	_			_	_				_				-		2736
	cga Arg	_	-	-			_	_	-							2784
_	cag Gln 930		_		_			-	-		_	_	_	-		2832
	tgg Trp															2880
	gag Glu															2928
	aga Arg															2976
								Ala					o A:		tg gac eu Asp	3024
	gac Asp 1010	Le					L G	aa ga lu Gi			ly L					3069
_	ggt Gly 1025	Sei					. P	ct ac ro Ti			hr L		acc o			3114
_	ggg Gly 1040	Sei		_			s د	gt co er Pi			er G					3159

-	aac Asn 1055	cag Gln	agc Ser	aac Asn	ctt Leu	ggg Gly 1060	gag Glu	gct Ala	tgt Cys	ctg Leu	gat Asp 1065	tct Ser	gcg Ala		3204
-	ggg Gly 1070					ttc Phe 1075									3249
						acg Thr 1090									3294
_	ggc Gly 1100			_	_	ctc Leu 1105			_	-		_	-		3339
_		-		_		agc Ser 1120				-		-	_	-	3384
	cat His 1130	_				agc Ser 1135					gtc Val 1140		ccg Pro		3429
	cca Pro 1145					gaa Glu 1150									3474
	gat Asp 1160	_			_	ggt Gly 1165					cgg Arg 1170	_			3519
	tcg Ser 1175					agt Ser 1180					acc Thr 1185		gag Glu		3564
_	gaa Glu 1190					gaa Glu 1195		_			aag Lys 1200		agg Arg		3609
						aga Arg 1210									3654
						ggt Gly 1225									3699
_	acg Thr 1235					ctc Leu 1240									3744
						gag Glu 1255									3789
						gga Gly 1270									3834

-		_	_	-		ggg Gly 1285		_	_	_	Āla		_	3879
					_	ccc Pro 1300	_	_		_	_			3924
	-	-	-		-	gcc Ala 1315	_		-		_			3969
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Leu Asn Gly Leu Ser Val Thr Gly Asp Ala Asp Asn Gln Tyr Gln Thr 35 40 45

Leu Tyr Lys Leu Tyr Glu Lys Cys Glu Val Val Met Gly Asn Leu Glu 50 55 60

Ile Val Leu Thr Gly His Asn Ala Asp Leu Ser Phe Leu Gln Trp Ile 70 75 80

Arg Glu Val Thr Gly Tyr Val Leu Val Ala Met Asn Glu Phe Ser Val 85 90 95

Leu Pro Leu Pro Asn Leu Arg Val Val Arg Gly Thr Gln Val Tyr Asp 100 105 110

Gly Lys Phe Ala Ile Phe Val Met Leu Asn Tyr Asn Thr Asn Ser Ser 115 120 125

His	Ala 130	Leu	Arg	Gln	Leu	Arg 135	Phe	Thr	Gln	Leu	Thr 140	Glu	Ile	Leu	Leu
Gly 145	Gly	Val	Tyr	Ile	Glu 150	Lys	Asn	Asp	Lys	Leu 155	Cys	His	Met	Asp	Thr 160
Ile	Asp	Trp	Arg	Asp 165	Ile	Val	Arg	Val	Pro 170	Asp	Ala	Glu	Ile	Val 175	Val
Lys	Asn	Asn	Gly 180	Gly	Asn	Cys	Pro	Pro 185	Cys	His	Glu	Val	Cys 190	Lys	Gly
Arg	Cys	Trp 195	Gly	Pro	Gly	Pro	Glu 200	Asp	Cys	Gln	Ile	Leu 205	Thr	Lys	Thr
Ile	Cys 210	Ala	Pro	Gln	Cys	Asn 215	Gly	Arg	Cys	Phe	Gly 220	Pro	Asn	Pro	Asn
Gln 225	Cys	Cys	His	Asp	Glu 230	Cys	Ala	Gly	Gly	Cys 235	Ser	Gly	Pro	Gln	Asp 240
Thr	Asp	Cys	Phe	Ala 245	Cys	Arg	His	Phe	Asn 250	Asp	Ser	Gly	Ala	Cys 255	Val
Pro	Arg	Суѕ	Pro 260	Ala	Pro	Leu	Val	Tyr 265	Asn	Lys	Leu	Thr	Phe 270	Gln	Leu
Glu	Pro	Asn 275	Pro	His	Ile	Lys	Tyr 280	Gln	Tyr	Gly	Gly	Val 285	Cys	Val	Ala
Ser	Cys 290		His	Asn	Phe			Asp				_	Val	Arg	Ala
Cys 305	Pro	Ala	Asp	Lys	Met 310	Glu	Val	Asp	Lys	Asn 315	Gly	Leu	Lys	Met	Cys 320
Glu	Pro	Cys	Arg	Gly 325	Leu	Cys	Pro	Lys	Ala 330	Cys	Glu	Gly	Thr	Gly 335	Ser
Gly	Ser	Arg	Tyr 340	Gln	Thr	Val	Asp	Ser 345	Ser	Asn	Ile	Asp	Gly 350	Phe	Val
Asn	Cys	Thr 355	Lys	Ile	Leu	Gly	Asn 360	Leu	Asp	Phe	Leu	Ile 365	Thr	Gly	Leu

Asn Gly Asp Pro Trp His Lys Ile Pro Ala Leu Asp Pro Glu Lys Leu Asn Val Phe Arg Thr Val Arg Glu Ile Thr Gly Tyr Leu Asn Ile Gln Ser Trp Pro Pro His Met His Asn Phe Ser Val Phe Ser Asn Leu Thr 405 410 Thr Ile Gly Gly Arg Ser Leu Tyr Asn Arg Gly Phe Ser Leu Leu Ile Met Lys Asn Leu Asn Val Thr Ser Leu Gly Phe Arg Ser Leu Lys Glu 440 Ile Ser Ala Gly Arg Val Tyr Ile Ser Ala Asn Gln Gln Leu Cys Tyr His His Ser Leu Asn Trp Thr Arg Leu Leu Arg Gly Pro Ala Glu Glu Arg Leu Asp Ile Lys Tyr Asn Arg Pro Leu Gly Glu Cys Val Ala Glu 490 Gly Lys Val Cys Asp Pro Leu Cys Ser Ser Gly Gly Cys Trp Gly Pro 500 505 Gly Pro Gly Gln Cys Leu Ser Cys Arg Asn Tyr Ser Arg Glu Gly Val 520 515 Cys Val Thr His Cys Asn Val Leu Gln Gly Glu Pro Arg Glu Phe Val 530 535 540 His Glu Ala His Cys Phe Ser Cys His Pro Glu Cys Gln Pro Met Glu 545 550 555 Gly Thr Ser Thr Cys Asn Gly Ser Gly Ser Asp Ala Cys Ala Arg Cys 565 570 Ala His Phe Arg Asp Gly Pro His Cys Val Asn Ser Cys Pro His Gly

Ile Leu Gly Ala Lys Gly Pro Ile Tyr Lys Tyr Pro Asp Ala Gln Asn

Glu Cys Arg Pro Cys His Glu Asn Cys Thr Gln Gly Cys Lys Gly Pro Glu Leu Gln Asp Cys Leu Gly Gln Ala Glu Val Leu Met Ser Lys Pro His Leu Val Ile Ala Val Thr Val Gly Leu Thr Val Ile Phe Leu Ile Leu Gly Gly Ser Phe Leu Tyr Trp Arg Gly Arg Arg Ile Gln Asn Lys Arg Ala Met Arg Arg Tyr Leu Glu Arg Gly Glu Ser Ile Glu Pro Leu Asp Pro Ser Glu Lys Ala Asn Lys Val Leu Ala Arg Ile Phe Lys Glu Thr Glu Leu Arg Lys Leu Lys Val Leu Gly Ser Gly Val Phe Gly Thr Val His Lys Gly Ile Trp Ile Pro Glu Gly Glu Ser Ile Lys Ile Pro Val Cys Ile Lys Val Ile Glu Asp Lys Ser Gly Arg Gln Ser Phe Gln Ala Val Thr Asp His Met Leu Ala Val Gly Ser Leu Asp His Ala His Ile Val Arg Leu Leu Gly Leu Cys Pro Gly Ser Ser Leu Gln Leu Val Thr Gln Tyr Leu Pro Leu Gly Ser Leu Leu Asp His Val Arg Gln His Arg Glu Thr Leu Gly Pro Gln Leu Leu Asn Trp Gly Val Gln Ile Ala Lys Gly Met Tyr Tyr Leu Glu Glu His Ser Met Val His Arg Asp Leu Ala Leu Arg Asn Val Met Leu Lys Ser Pro Ser Gln Val Gln Val

- Ala Asp Phe Gly Val Ala Asp Leu Leu Pro Pro Asp Asp Lys Gln Leu 850 855 860
- Leu His Ser Glu Ala Lys Thr Pro Ile Lys Trp Met Ala Leu Glu Ser 865 870 875 880
- Ile His Phe Gly Lys Tyr Thr His Gln Ser Asp Val Trp Ser Tyr Gly 885 890 895
- Val Thr Val Trp Glu Leu Met Thr Phe Gly Ala Glu Pro Tyr Ala Gly 900 905 910
- Leu Arg Leu Ala Glu Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg Leu 915 920 925
- Ala Gln Pro Gln Ile Cys Thr Ile Asp Val Tyr Met Val Met Val Lys 930 935 940
- Cys Trp Met Ile Asp Glu Asn Ile Arg Pro Thr Phe Lys Glu Leu Ala 945 950 955 960
- Asn Glu Phe Thr Arg Met Ala Arg Asp Pro Pro Arg Tyr Leu Val Ile 965 970 975
- Lys Arg Ala Ser Gly Pro Gly Ile Pro Pro Ala Ala Glu Pro Ser Ala 980 985 990
- Leu Ser Thr Lys Glu Leu Gln Asp Ala Glu Leu Glu Pro Asp Leu Asp 995 1000 1005
- Leu Asp Leu Asp Val Glu Val Glu Glu Glu Gly Leu Ala Thr Thr 1010 1015 1020
- Leu Gly Ser Ala Leu Ser Leu Pro Thr Gly Thr Leu Thr Arg Pro 1025 1030 1035
- Arg Gly Ser Gln Ser Leu Leu Ser Pro Ser Ser Gly Tyr Met Pro 1040 1045 1050
- Met Asn Gln Ser Asn Leu Gly Glu Ala Cys Leu Asp Ser Ala Val 1055 1060 1065
- Leu Gly Gly Arg Glu Gln Phe Ser Arg Pro Ile Ser Leu His Pro 1070 1075 1080

- Ile Pro Arg Gly Arg Gln Thr Ser Glu Ser Ser Glu Gly His Val 1085 1090 1095
- Thr Gly Ser Glu Ala Glu Leu Gln Glu Arg Val Ser Met Cys Arg 1100 1105 1110
- Ser Arg Ser Arg Ser Arg Ser Pro Arg Pro Arg Gly Asp Ser Ala 1115 1120 1125
- Tyr His Ser Gln Arg His Ser Leu Leu Thr Pro Val Thr Pro Leu 1130 1135 1140
- Ser Pro Pro Gly Leu Glu Glu Glu Asp Gly Asn Gly Tyr Val Met 1145 1150 1155
- Pro Asp Thr His Leu Arg Gly Thr Ser Ser Ser Arg Glu Gly Thr 1160 1165 1170
- Leu Ser Ser Val Gly Leu Ser Ser Val Leu Gly Thr Glu Glu Glu 1175 1180 1185
- Asp Glu Asp Glu Glu Tyr Glu Tyr Met Asn Arg Lys Arg Arg Gly 1190 1200
- Ser Pro Ala Arg Pro Pro Arg Pro Gly Ser Leu Glu Glu Leu Gly 1205 1210 1215
- Tyr Glu Tyr Met Asp Val Gly Ser Asp Leu Ser Ala Ser Leu Gly 1220 1235
- Ser Thr Gln Ser Cys Pro Leu His Pro Met Ala Ile Val Pro Ser 1235 1240 1245
- Ala Gly Thr Thr Pro Asp Glu Asp Tyr Glu Tyr Met Asn Arg Arg 1250 1260
- Arg Gly Ala Gly Gly Ser Gly Gly Asp Tyr Ala Ala Met Gly Ala 1265 1270 1275
- Cys Pro Ala Ala Glu Gln Gly Tyr Glu Glu Met Arg Ala Phe Gln 1280 1285 1290
- Gly Pro Gly His Gln Ala Pro His Val Arg Tyr Ala Arg Leu Lys 1295 1300 1305

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Thr Leu Arg Ser Leu Glu Ala Thr Asp Ser Ala Phe Asp Asn Pro

1315

1310

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	ttc Phe				-				_		_	534
	cca Pro 170											582
	tgt Cys											630
	gaa Glu											678
	gac Asp											726
	tgt Cys											774
	atg Met 250											822
	ttt Phe											870
	aag Lys											918
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	gaa Glu											1014
	tgc Cys 330											1062
	cag Gln											1110
	atc Ile											1158

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					gag Glu										1254
					gac Asp										1302
		_	_		tat Tyr	_		_		_			_		1350
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					act Thr	_		_		_	_				1446
					ctc Leu		_				_	_	_		1494
					gct Ala										1542
					agt Ser										1590
					cgc Arg 525										1638
					ggt Gly										1686
					ccc Pro										1734
aca .Thr	_			_	ggt Gly		-		_		-	_			1782
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					gag Glu											22	14
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					ttc Phe 765											23	58
					gtc Val											24	06
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gtc Val	cac His	gag Glu 810	cac His	aag Lys	gat Asp	aac Asn	att Ile 815	gga Gly	tca Ser	caa Gln	ctg Leu	ctg Leu 820	ctt Leu	aac Asn	tgg Trp	25	02
					aag Lys											25	50
gtt Val 840	cat His	cgg Arg	gat Asp	ttg Leu	gca Ala 845	gcc Ala	cgt Arg	aat Asn	gtc Val	tta Leu 850	gtg Val	aaa Lys	tct Ser	cca Pro	aac Asn 855	2.5	98

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gct ctg gag tgt ata cat tac agg aaa ttc acc cat cag agt gac gtt Ala Leu Glu Cys Ile His Tyr Arg Lys Phe Thr His Gln Ser Asp Val 890 895 900	
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ccc tat gat gga att cca acg cga gaa atc cct gat tta tta gag aaa Pro Tyr Asp Gly Ile Pro Thr Arg Glu Ile Pro Asp Leu Leu Glu Lys 920 925 930 935	
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caa Gln 1120					acc Thr 1125										3435
ttt Phe 1135					agc Ser 1140										3480
tac Tyr 1150	Met				cga Arg 1155										3525
cca Pro 1165					cct Pro 1170										3570
ctt Leu 1180					aat Asn 1185										3615
cca Pro 1195					gat Asp 1200								tac Tyr		3660
aac Asn 1210					acc Thr 1215							_	aag Lys		3705
aac Asn 1225					cca Pro 1230								gac Asp		3750
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Glu Asn Lys Leu Ser Ser Leu Ser Asp Leu Glu Gln Gln Tyr Arg Ala 35 40 45

Leu Arg Lys Tyr Tyr Glu Asn Cys Glu Val Val Met Gly Asn Leu Glu 50 55 60

Ile Thr Ser Ile Glu His Asn Arg Asp Leu Ser Phe Leu Arg Ser Val 65 70 75 80

Arg Glu Val Thr Gly Tyr Val Leu Val Ala Leu Asn Gln Phe Arg Tyr 85 90 95

Leu Pro Leu Glu Asn Leu Arg Ile Ile Arg Gly Thr Lys Leu Tyr Glu
100 105 110

Asp Arg Tyr Ala Leu Ala Ile Phe Leu Asn Tyr Arg Lys Asp Gly Asn 115 120 125

Phe Gly Leu Gln Glu Leu Gly Leu Lys Asn Leu Thr Glu Ile Leu Asn 130 135 140

Gly Gly Val Tyr Val Asp Gln Asn Lys Phe Leu Cys Tyr Ala Asp Thr 145 150 155 160

Ile His Trp Gln Asp Ile Val Arg Asn Pro Trp Pro Ser Asn Leu Thr
165 170 175

Leu Val Ser Thr Asn Gly Ser Ser Gly Cys Gly Arg Cys His Lys Ser 180 185 190

Cys Thr Gly Arg Cys Trp Gly Pro Thr Glu Asn His Cys Gln Thr Leu 195 200 205

Thr Arg Thr Val Cys Ala Glu Gln Cys Asp Gly Arg Cys Tyr Gly Pro 210 215 220

Tyr Val Ser Asp Cys Cys His Arg Glu Cys Ala Gly Gly Cys Ser Gly 225 230 235 240

Pro Lys Asp Thr Asp Cys Phe Ala Cys Met Asn Phe Asn Asp Ser Gly 245 250 255

Ala Cys Val Thr Gln Cys Pro Gln Thr Phe Val Tyr Asn Pro Thr Thr 260 265 270

Phe Gln Leu Glu His Asn Phe Asn Ala Lys Tyr Thr Tyr Gly Ala Phe 275 280 285

Cys Val Lys Lys Cys Pro His Asn Phe Val Val Asp Ser Ser Ser Cys 290 295 300

Val Arg Ala Cys Pro Ser Ser Lys Met Glu Val Glu Glu Asn Gly Ile 305 310 315 320

Lys Met Cys Lys Pro Cys Thr Asp Ile Cys Pro Lys Ala Cys Asp Gly 325 330 335

Ile Gly Thr Gly Ser Leu Met Ser Ala Gln Thr Val Asp Ser Ser Asn 340 345 350

Ile Asp Lys Phe Ile Asn Cys Thr Lys Ile Asn Gly Asn Leu Ile Phe 355 360 365

Leu Val Thr Gly Ile His Gly Asp Pro Tyr Asn Ala Ile Glu Ala Ile 370 375 380

Asp Pro Glu Lys Leu Asn Val Phe Arg Thr Val Arg Glu Ile Thr Gly 385 390 395

Phe Leu Asn Ile Gln Ser Trp Pro Pro Asn Met Thr Asp Phe Ser Val405 410 415

Phe Ser Asn Leu Val Thr Ile Gly Gly Arg Val Leu Tyr Ser Gly Leu 420 425 430

Ser Leu Leu Ile Leu Lys Gln Gln Gly Ile Thr Ser Leu Gln Phe Gln 435 440 445

Ser Leu Lys Glu Ile Ser Ala Gly Asn Ile Tyr Ile Thr Asp Asn Ser 450 455 460

Asn Leu Cys Tyr Tyr His Thr Ile Asn Trp Thr Thr Leu Phe Ser Thr 465 470 475 480

. . . .

Ile Asn Gln Arg Ile Val Ile Arg Asp Asn Arg Lys Ala Glu Asn Cys 485 490 495

Thr Ala Glu Gly Met Val Cys Asn His Leu Cys Ser Ser Asp Gly Cys 500 505 510

Trp Gly Pro Gly Pro Asp Gln Cys Leu Ser Cys Arg Arg Phe Ser Arg 515 520 525

Gly Arg Ile Cys Ile Glu Ser Cys Asn Leu Tyr Asp Gly Glu Phe Arg 530 535 540

Glu Phe Glu Asn Gly Ser Ile Cys Val Glu Cys Asp Pro Gln Cys Glu 545 550 555 560

Lys Met Glu Asp Gly Leu Leu Thr Cys His Gly Pro Gly Pro Asp Asn 565 570 575

Cys Thr Lys Cys Ser His Phe Lys Asp Gly Pro Asn Cys Val Glu Lys 580 585 590

Cys Pro Asp Gly Leu Gln Gly Ala Asn Ser Phe Ile Phe Lys Tyr Ala 595 600 605

Asp Pro Asp Arg Glu Cys His Pro Cys His Pro Asn Cys Thr Gln Gly 610 615 620

Cys Asn Gly Pro Thr Ser His Asp Cys Ile Tyr Tyr Pro Trp Thr Gly 625 630 635 640

His Ser Thr Leu Pro Gln His Ala Arg Thr Pro Leu Ile Ala Ala Gly 645 650 655

Val Ile Gly Gly Leu Phe Ile Leu Val Ile Val Gly Leu Thr Phe Ala 660 665 670

Val Tyr Val Arg Arg Lys Ser Ile Lys Lys Lys Arg Ala Leu Arg Arg 675 680 685

Phe Leu Glu Thr Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Thr Ala 690 695 700 Pro Asn Gln Ala Gln Leu Arg Ile Leu Lys Glu Thr Glu Leu Lys Arg 705 710 715 720

Val Lys Val Leu Gly Ser Gly Ala Phe Gly Thr Val Tyr Lys Gly Ile
725 730 735

Trp Val Pro Glu Gly Glu Thr Val Lys Ile Pro Val Ala Ile Lys Ile 740 745 750

Leu Asn Glu Thr Thr Gly Pro Lys Ala Asn Val Glu Phe Met Asp Glu 755 760 765

Ala Leu Ile Met Ala Ser Met Asp His Pro His Leu Val Arg Leu Leu 770 775 780

Gly Val Cys Leu Ser Pro Thr Ile Gln Leu Val Thr Gln Leu Met Pro 785 790 795 800

His Gly Cys Leu Leu Glu Tyr Val His Glu His Lys Asp Asn Ile Gly 805 810 815

Ser Gln Leu Leu Asn Trp Cys Val Gln Ile Ala Lys Gly Met Met 820 825 830

Tyr Leu Glu Glu Arg Arg Leu Val His Arg Asp Leu Ala Ala Arg Asn 835 840 845

Val Leu Val Lys Ser Pro Asn His Val Lys Ile Thr Asp Phe Gly Leu 850 855 860

Ala Arg Leu Leu Glu Gly Asp Glu Lys Glu Tyr Asn Ala Asp Gly Gly 865 870 875 880

Lys Met Pro Ile Lys Trp Met Ala Leu Glu Cys Ile His Tyr Arg Lys 885 890 895

Phe Thr His Gln Ser Asp Val Trp Ser Tyr Gly Val Thr Ile Trp Glu 900 905 910

Leu Met Thr Phe Gly Gly Lys Pro Tyr Asp Gly Ile Pro Thr Arg Glu 915 920 925

Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg Leu Pro Gln Pro Pro Ile 930 935 940

. . .

- Cys Thr Ile Asp Val Tyr Met Val Met Val Lys Cys Trp Met Ile Asp 945 950 955 960
- Ala Asp Ser Arg Pro Lys Phe Lys Glu Leu Ala Ala Glu Phe Ser Arg 965 970 975
- Met Ala Arg Asp Pro Gln Arg Tyr Leu Val Ile Gln Gly Asp Asp Arg 980 985 990
- Met Lys Leu Pro Ser Pro Asn Asp Ser Lys Phe Phe Gln Asn Leu Leu 995 1000 1005
- Asp Glu Glu Asp Leu Glu Asp Met Met Asp Ala Glu Glu Tyr Leu 1010 1015 1020
- Val Pro Gln Ala Phe Asn Ile Pro Pro Pro Ile Tyr Thr Ser Arg 1025 1030 1035
- Ala Arg Ile Asp Ser Asn Arg Ser Glu Ile Gly His Ser Pro Pro 1040 1045 1050
- Pro Ala Tyr Thr Pro Met Ser Gly Asn Gln Phe Val Tyr Arg Asp 1055 1060 1065
- Gly Gly Phe Ala Ala Glu Gln Gly Val Ser Val Pro Tyr Arg Ala 1070 1075 1080
- Pro Thr Ser Thr Ile Pro Glu Ala Pro Val Ala Gln Gly Ala Thr 1085 1090 1095
- Pro Val Ala Pro His Val Gln Glu Asp Ser Ser Thr Gln Arg Tyr 1115 1120 1125
- Ser Ala Asp Pro Thr Val Phe Ala Pro Glu Arg Ser Pro Arg Gly 1130 1135 1140
- Glu Leu Asp Glu Glu Gly Tyr Met Thr Pro Met Arg Asp Lys Pro 1145 1150 1155
- Lys Gln Glu Tyr Leu Asn Pro Val Glu Glu Asn Pro Phe Val Ser 1160 1165 1170

Arg	Arg 1175	Lys	Asn	Gly	Asp	Leu 1180		Ala	Leu	Asp	Asn 1185	Pro	Glu	Tyr	
	Asn 1190	Ala	Ser	Asn	Gly	Pro 1195		Lys	Ala	Glu	Asp 1200	Glu	Tyr	Val	
Asn	Glu 1205	Pro	Leu	Tyr	Leu	Asn 1210		Phe	Ala	Asn	Thr 1215	Leu	Gly	Lys	
Ala	Glu 1220	Tyr	Leu	Lys	Asn	Asn 1225		Leu	Ser	Met	Pro 1230	Glu	Lys	Ala	
Lys	Lys 1235	Ala	Phe	Asp	Asn	Pro 1240		Tyr	Trp	Asn	His 1245	Ser	Leu	Pro	
Pro	Arg 1250	Ser	Thr	Leu	Gln	His 1255	Pro	Asp	Tyr	Leu	Gln 1260	Glu	Tyr	Ser	
Thr	Lys 1265	Tyr	Phe	Tyr	Lys	Gln 1270		Gly	Arg	Ile	Arg 1275	Pro	Ile	Val	
Ala	Glu 1280	Asn	Pro	Glu	Tyr	Leu 1285	Ser	Glu	Phe	Ser	Leu 1290	Lys	Pro	Gly	
Thr	Val 1295	Leu	Pro	Pro	Pro	Pro 1300		Arg	His	Arg	Asn 1305	Thr	Val	Val	
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<400 atg Met 1	gca a	ata d	Pro S	ca a Ser <i>l</i>	aga d Arg <i>P</i>	egc aa Arg Ly	ag ga ys As	ac ta sp Ty 10	r Pı	ca aa co Ly	aa cta /s Leu	ı tct ı Ser	gaa Glu 15	atc Ile	48
		Cys A		_	-	-		s Le		-	et cto er Leu		_	_	96
	Gln (rg Ly				aa aac Lu Asr 45				144

				agc Ser						:	192
				gtc Val						2	240
				ctg Leu						2	288
				tat Tyr 105						;	336
				ctc Leu						Ş	384
_	_			 gtc Val	_	_	_			4	432
				tgg Trp						4	480
		_	_	 tca Ser			_		-		528
				ggc Gly 185							576
				act Thr						•	624
				agt Ser						•	672
				gac Asp						-	720
				gtt Val						Ī	768
				ctg Leu 265						8	316
				aag Lys						8	364

					_	gta Val 295	_	_	_		_		_	_	-		912
						aaa Lys											960
						atc Ile]	1008
						att Ile										1	1056
						ctt Leu										1	1104
						gat Asp 375										1	1152
						ttc Phe										1	1200
						ttc Phe										1	1248
						tca Ser	_	_								1	1296
						tct Ser										1	L344
						aac Asn 455										1	1392
aca Thr 465	aca Thr	ctc Leu	ttc Phe	agc Ser	acc Thr 470	att Ile	aac Asn	cag Gln	aga Arg	ata Ile 475	gtg Val	atc Ile	cga Arg	gat Asp	aac Asn 480	1	440
aga Arg	aga Arg	gct Ala	gag Glu	aat Asn 485	tgt Cys	act Thr	gct Ala	gaa Glu	ggc Gly 490	atg Met	gta Val	tgc Cys	aac Asn	cac His 495	ctg Leu	1	.488
tgt Cys	tca Ser	aat Asn	gat Asp 500	ggt Gly	tgt Cys	tgg Trp	gga Gly	cct Pro 505	ggg Gly	ccg Pro	gac Asp	cag Gln	tgc Cys 510	ctg Leu	tca Ser	1	.536
						gga Gly										1	.584

. 4 . tat gat ggg gaa ttt cga gag ttt gaa aac ggc tcc atc tgt gtt gag 1632 Tyr Asp Gly Glu Phe Arg Glu Phe Glu Asn Gly Ser Ile Cys Val Glu 535 tgt gac tcc cag tgt gag aaa atg gaa gat gga ctc ctc aca tgc cat 1680 Cys Asp Ser Gln Cys Glu Lys Met Glu Asp Gly Leu Leu Thr Cys His 550 gga ccg gga cct gac aac tgc aca aag tgc tct cat ttt aag gat ggt 1728 Gly Pro Gly Pro Asp Asn Cys Thr Lys Cys Ser His Phe Lys Asp Gly 570 cca aac tgt gtg gag aaa tgt cca gat ggc cta cag gga gca aac agt 1776 Pro Asn Cys Val Glu Lys Cys Pro Asp Gly Leu Gln Gly Ala Asn Ser 585 ttc att ttt aag tat gca gat cag gat cgg gag tgc cac cct tgc cat 1824 Phe Ile Phe Lys Tyr Ala Asp Gln Asp Arg Glu Cys His Pro Cys His 600 cca aac tgc acc cag ggg tgt aac ggt ccc act agt cat gac tgc att 1872 Pro Asn Cys Thr Gln Gly Cys Asn Gly Pro Thr Ser His Asp Cys Ile tac tac cca tgg acg ggc cat tcc act tta cca caa cac gct aga act 1920 Tyr Tyr Pro Trp Thr Gly His Ser Thr Leu Pro Gln His Ala Arg Thr 630 635 1968 cca ctg att gca gcc gga gtc att gga ggc ctc ttc atc ctg gtg atc Pro Leu Ile Ala Ala Gly Val Ile Gly Gly Leu Phe Ile Leu Val Ile 650 atg gct ttg aca ttt gct gtc tat gtc aga aga aag agc atc aaa aag 2016 Met Ala Leu Thr Phe Ala Val Tyr Val Arg Arg Lys Ser Ile Lys Lys aaa cgt gct ttg agg aga ttc ctg gag aca gag ctg gta gag ccc tta 2064 Lys Arg Ala Leu Arg Arg Phe Leu Glu Thr Glu Leu Val Glu Pro Leu 680 685 act ccc agt ggc acg gca ccc aat caa gct caa ctt cgc att ttg aag 2112 Thr Pro Ser Gly Thr Ala Pro Asn Gln Ala Gln Leu Arg Ile Leu Lys 690 695 700 gaa acc gaa cta aag agg gta aag gtc ctt ggc tcg gga gct ttt gga 2160 Glu Thr Glu Leu Lys Arg Val Lys Val Leu Gly Ser Gly Ala Phe Gly 705 715 acc gtt tat aaa ggt att tgg gtg cct gaa ggt gaa aca gtg aaa atc 2208 Thr Val Tyr Lys Gly Ile Trp Val Pro Glu Gly Glu Thr Val Lys Ile 725 730

cct gtg gct ata aag atc ctc aat gaa aca act ggc ccc aaa gcc aac

Pro Val Ala Ile Lys Ile Leu Asn Glu Thr Thr Gly Pro Lys Ala Asn

2256

2304

		_	_		_	gga Gly 775		_	_	_				_	-	2352
-	_	_	_	_	_	cat His		_		_	-		_		•	2400
	_	-				tca Ser	_	_	_	_			-	_	-	2448
						tac Tyr		_	_	-			_			2496
						gtc Val										2544
atc Ile	aca Thr 850	gat Asp	ttt Phe	gga Gly	ctg Leu	gcc Ala 855	cgg Arg	ctc Leu	ttg Leu	gaa Glu	gga Gly 860	gat Asp	gaa Glu	aaa Lys	gaa Glu	2592
						aag Lys										2640
						ttc Phe										2688
						ctg Leu										2736
						atc Ile										2784
						tgc Cys 935										2832
						gct Ala										2880
						atg Met										2928
						atg Met										2976
						gat Asp		Glu	g gat 1 Asp) Me	-	g gat et Asp	3024

 gag Glu 1010	_	_	_	ccc Pro 1015	_	_						3069
 tac Tyr 1025				aga Arg 1030				agg Arg 1035				3114
cac His 1040				gcc Ala 1045								3159
gtg Val 1055				ggc Gly 1060								3204
				acc Thr 1075								3249
				gag Glu 1090								3294
acc Thr 1100				gtg Val 1105								3339
				gct Ala 1120								3384
 aat Asn 1130				ctg Leu 1135						act Thr		3429
				caa Gln 1150								3474
cct Pro 1160				agg Arg 1165							tta Leu	3519
aat Asn 1175				agt Ser 1180						aag Lys		3564
gat Asp 1190				gag Glu 1195						ttc Phe	_	3609
gcc Ala 1205	_	 _	_	gag Glu 1210		_		_	_	_		3654
 cca Pro 1220				aaa Lys 1225								3699

aac cac agc ctg cca ccc cgg agc acc ctt cag cac cca gac tac Asn His Ser Leu Pro Pro Arg Ser Thr Leu Gln His Pro Asp Tyr 1235 1240 1245	3744
ctg cag gaa tac agc aca aaa tat ttt tat aaa cag aat gga cgg Leu Gln Glu Tyr Ser Thr Lys Tyr Phe Tyr Lys Gln Asn Gly Arg 1250 1255 1260	3789
atc cgc ccc att gtg gca gag aat cct gag tac ctc tcg gag ttc Ile Arg Pro Ile Val Ala Glu Asn Pro Glu Tyr Leu Ser Glu Phe 1265 1270 1275	3834
tcg ctg aag cct ggc act atg ctg ccc cct ccg ccc tac aga cac Ser Leu Lys Pro Gly Thr Met Leu Pro Pro Pro Pro Tyr Arg His 1280 1285 1290	3879
cgg aat act gtg gtg tga Arg Asn Thr Val Val 1295	3897
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Thr Val Cys Ala Gly Thr Glu Asn Lys Leu Ser Ser Leu Ser Asp Leu 20 25 30	
Glu Gln Gln Tyr Arg Ala Leu Arg Lys Tyr Tyr Glu Asn Cys Glu Val 35 40 45	
Val Met Gly Asn Leu Glu Ile Thr Ser Ile Glu His Asn Arg Asp Leu 50 55 60	
Ser Phe Leu Arg Ser Ile Arg Glu Val Thr Gly Tyr Val Leu Val Ala 65 70 75 80	
Leu Asn Gln Phe Arg Tyr Leu Pro Leu Glu Asn Leu Arg Ile Ile Arg 85 90 95	
Gly Thr Lys Leu Tyr Glu Asp Arg Tyr Ala Leu Ala Ile Phe Leu Asn 100 105 110	
Tyr Arg Lys Asp Gly Asn Phe Gly Leu Gln Glu Leu Gly Leu Lys Asn 115 120 125	

. * 4) *

Leu Thr Glu Ile Leu Asn Gly Gly Val Tyr Val Asp Gln Asn Lys Phe Leu Cys Tyr Ala Asp Thr Ile His Trp Gln Asp Ile Val Arg Asn Pro Trp Pro Ser Asn Met Thr Leu Val Ser Thr Asn Gly Ser Ser Gly Cys 165 170 Gly Arg Cys His Lys Ser Cys Thr Gly Arg Cys Trp Gly Pro Thr Glu Asn His Cys Gln Thr Leu Thr Arg Thr Val Cys Ala Glu Gln Cys Asp Gly Arg Cys Tyr Gly Pro Tyr Val Ser Asp Cys Cys His Arg Glu Cys Ala Gly Gly Cys Ser Gly Pro Lys Asp Thr Asp Cys Phe Ala Cys Met 235 Asn Phe Asn Asp Ser Gly Ala Cys Val Thr Gln Cys Pro Gln Thr Phe 245 250 Val Tyr Asn Pro Thr Thr Phe Gln Leu Glu His Asn Phe Asn Ala Lys 260 265 270 Tyr Thr Tyr Gly Ala Phe Cys Val Lys Lys Cys Pro His Asn Phe Val 275 280 285 Val Asp Ser Ser Cys Val Arg Ala Cys Pro Ser Ser Lys Met Glu 290 295 300 Val Glu Glu Asn Gly Ile Lys Met Cys Lys Pro Cys Thr Asp Ile Cys 305 310 315 Pro Lys Ala Cys Asp Gly Ile Gly Thr Gly Ser Leu Met Ser Ala Gln 325 330 Thr Val Asp Ser Ser Asn Ile Asp Lys Phe Ile Asn Cys Thr Lys Ile 345 Asn Gly Asn Leu Ile Phe Leu Val Thr Gly Ile His Gly Asp Pro Tyr 360

- Asn Ala Ile Asp Ala Ile Asp Pro Glu Lys Leu Asn Val Phe Arg Thr $370 \hspace{1cm} 375 \hspace{1cm} 380$
- Val Arg Glu Ile Thr Gly Phe Leu Asn Ile Gln Ser Trp Pro Pro Asn 385 390 395 400
- Met Thr Asp Phe Ser Val Phe Ser Asn Leu Val Thr Ile Gly Gly Arg
 405 410 415
- Val Leu Tyr Ser Gly Leu Ser Leu Leu Ile Leu Lys Gln Gln Gly Ile 420 425 430
- Thr Ser Leu Gln Phe Gln Ser Leu Lys Glu Ile Ser Ala Gly Asn Ile 435 440 445
- Tyr Ile Thr Asp Asn Ser Asn Leu Cys Tyr Tyr His Thr Ile Asn Trp 450 455 460
- Thr Thr Leu Phe Ser Thr Ile Asn Gln Arg Ile Val Ile Arg Asp Asn 465 470 475 480
- Arg Arg Ala Glu Asn Cys Thr Ala Glu Gly Met Val Cys Asn His Leu 485 490 495
- Cys Ser Asn Asp Gly Cys Trp Gly Pro Gly Pro Asp Gln Cys Leu Ser 500 505 510
- Cys Arg Arg Phe Ser Arg Gly Lys Ile Cys Ile Glu Ser Cys Asn Leu 515 520 525
- Tyr Asp Gly Glu Phe Arg Glu Phe Glu Asn Gly Ser Ile Cys Val Glu 530 535 540
- Cys Asp Ser Gln Cys Glu Lys Met Glu Asp Gly Leu Leu Thr Cys His 545 550 . 555 560
- Gly Pro Gly Pro Asp Asn Cys Thr Lys Cys Ser His Phe Lys Asp Gly 565 570 575
- Pro Asn Cys Val Glu Lys Cys Pro Asp Gly Leu Gln Gly Ala Asn Ser 580 585 590
- Phe Ile Phe Lys Tyr Ala Asp Gln Asp Arg Glu Cys His Pro Cys His 595 600 605

Pro	Asn 610	Cys	Thr	Gln	Gly	Cys 615	Asn	Gly	Pro	Thr	Ser 620	His	Asp	Cys	Ile
Туг 625	Tyr	Pro	Trp	Thr	Gly 630	His	Ser	Thr	Leu	Pro 635	Gln	His	Ala	Arg	Thr 640
Pro	Leu	Ile	Ala	Ala 645	Gly	Val	Ile	Gly	Gly 650	Leu	Phe	Ile	Leu	Val 655	Ile
Met	Ala	Leu	Thr 660	Phe	Ala	Val	Tyr	Val 665	Arg	Arg	Lys	Ser	Ile 670	Lys	Lys
Lys	Arg	Ala 675	Leu	Arg	Arg	Phe	Leu 680	Glu	Thr	Glu	Leu	Val 685	Glu	Pro	Leu
Thr	Pro 690	Ser	Gly	Thr	Ala	Pro 695	Asn	Gln	Ala	Gln	Leu 700	Arg	Ile	Leu	Lys
Glu 705	Thr	Glu	Leu	Lys	Arg 710	Val	Lys	Val	Leu	Gly 715	Ser	Gly	Ala	Phe	Gly 720
Thr	Val	Tyr	Lys	Gly 725	Ile	Trp	Val	Pro	Glu 730	Gly	Glu	Thr	Val	Lys 735	Ile
Pro	Val	Ala	Ile 740	Lys	Ile	Leu	Asn	Glu 745	Thr	Thr	Gly	Pro	Lys 750	Ala	Asn
Val	Glu	Phe 755	Met	Asp	Glu	Ala	Leu 760	Ile	Met	Ala	Ser	Met 765	Asp	His	Pro
His	Leu 770	Val	Arg	Leu	Leu	Gly 775	Val	Cys	Leu	Ser	Pro 780	Thr	Ile	Gln	Leu
Val 785	Thr	Gln	Leu	Met	Pro 790	His	Gly	Cys	Leu	Leu 795	Asp	Tyr	Val	His	Glu 800
His	Lys	Asp	Asn	Ile 805	Gly	Ser	Gln	Leu	Leu 810	Leu	Asn	Trp	Cys	Val 815	Gln
Ile	Ala	Lys	Gly 820	Met	Met	Tyr	Leu	Glu 825	Glu	Arg	Arg	Leu	Val 830	His	Arg

Asp Leu Ala Ala Arg Asn Val Leu Val Lys Ser Pro Asn His Val Lys 835 840 845

- Ile Thr Asp Phe Gly Leu Ala Arg Leu Leu Glu Gly Asp Glu Lys Glu 850 855 860
- Tyr Asn Ala Asp Gly Gly Lys Met Pro Ile Lys Trp Met Ala Leu Glu . 865 870 875 880
- Cys Ile His Tyr Arg Lys Phe Thr His Gln Ser Asp Val Trp Ser Tyr 885 890 895
- Gly Val Thr Ile Trp Glu Leu Met Thr Phe Gly Gly Lys Pro Tyr Asp 900 905 910
- Gly Ile Pro Thr Arg Glu Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg 915 920 925
- Leu Pro Gln Pro Pro Ile Cys Thr Ile Asp Val Tyr Met Val Met Val 930 935 940
- Lys Cys Trp Met Ile Asp Ala Asp Ser Arg Pro Lys Phe Lys Glu Leu 945 950 955 960
- Ala Ala Glu Phe Ser Arg Met Ala Arg Asp Pro Gln Arg Tyr Leu Val $965 \hspace{1.5cm} 970 \hspace{1.5cm} 975$
- Ile Gln Gly Asp Asp Arg Met Lys Leu Pro Ser Pro Asn Asp Ser Lys 980 985 990
- Phe Phe Gln Asn Leu Leu Asp Glu Glu Asp Leu Glu Asp Met Met Asp 995 1000 1005
- Ala Glu Glu Tyr Leu Val Pro Gln Ala Phe Asn Ile Pro Pro Pro 1010 1015 1020
- Ile Tyr Thr Ser Arg Thr Arg Ilé Asp Ser Asn Arg Ser Glu Ile 1025 1030 1035
- Gly His Ser Pro Pro Pro Ala. Tyr Thr Pro Met Ser Gly Asn Gln 1040 1045 1050
- Phe Val Tyr Gln Asp Gly Gly Phe Ala Thr Gln Gln Gly Met Pro 1055 1060 1065
- Met Pro Tyr Arg Ala Thr Thr Ser Thr Ile Pro Glu Ala Pro Val 1070 1075 1080

Ala Gln Gly Ala Thr Ala Glu Met Phe Asp Asp Ser Cys Cys Asn 1085 1090 1095

Gly Thr Leu Arg Lys Pro Val Ala Pro His Val Gln Glu Asp Ser 1100 1105 1110

Ser Thr Gln Arg Tyr Ser Ala Asp Pro Thr Val Phe Ala Pro Glu 1115 1120 1125

Arg Asn Pro Arg Gly Glu Leu Asp Glu Glu Gly Tyr Met Thr Pro 1130 1135 1140

Met His Asp Lys Pro Lys Gln Glu Tyr Leu Asn Pro Val Glu Glu 1145 1150 1155

Asn Pro Phe Val Ser Arg Arg Lys Asn Gly Asp Leu Gln Ala Leu 1160 1165 1170

Asp Asn Pro Glu Tyr His Ser Ala Ser Ser Gly Pro Pro Lys Ala 1175 1180 1185

Glu Asp Glu Tyr Val Asn Glu Pro Leu Tyr Leu Asn Thr Phe Ala 1190 1200

Asn Ala Leu Gly Ser Ala Glu Tyr Met Lys Asn Ser Val Leu Ser 1205 1210 . 1215

Val Pro Glu Lys Ala Lys Lys Ala Phe Asp Asn Pro Asp Tyr Trp 1220 1225 1230

Asn His Ser Leu Pro Pro Arg Ser Thr Leu Gln His Pro Asp Tyr 1235 1240 1245

Leu Gln Glu Tyr Ser Thr Lys Tyr Phe Tyr Lys Gln Asn Gly Arg 1250 1260

Ile Arg Pro Ile Val Ala Glu Asn Pro Glu Tyr Leu Ser Glu Phe 1265 1270 1275

Ser Leu Lys Pro Gly Thr Met Leu Pro Pro Pro Pro Tyr Arg His 1280 1285 1290

Arg Asn Thr Val Val 1295